

SEQ ID NO: 1 OF  
5177,188

# AntiJen

a kinetic, thermodynamic and cellular database

## FULL SEARCH: T CELL EPITOPE

1 Enter an epitope string

- Replace variable amino acid positions with an asterisk (\*).
- Encase alternative amino acids in square brackets e.g. [AEI]

2 Enter the length of amino acid string

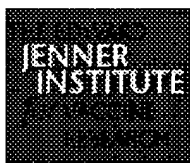
Leave blank to retrieve all.

MIN  MAX

3 Select the restriction allele or serotype

New Search: [MHC Ligand](#) [MHC Kinetics](#) [T Cell Epitope](#) [TCR-MHC](#) [TAP Ligand](#) [Protein I](#)  
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## THE ANTIJEN DATABASE SEARCH RESULTS

You searched for epitope string "YHDRKEFAKFEEERARAKWDTANN" of undefined length.

Sorry there are no results for your search criteria

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## FULL SEARCH: T CELL EPITOPE

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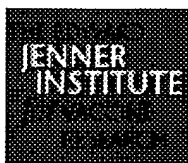
2 Enter the length of amino acid string  
Leave blank to retrieve all.

MIN  MAX

3 Select the restriction allele or serotype

New Search: [MHC Ligand](#) [MHC Kinetics](#) [T Cell Epitope](#) [TCR-MHC](#) [TAP Ligand](#) [Protein L](#)  
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## THE ANTIJEN DATABASE SEARCH RESULTS

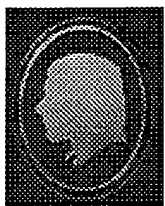
You searched for epitope string "YHDRKEFAKFEEERARAKWDTANN" of undefined length.

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## FULL SEARCH: T CELL EPITOPE

1 Enter an epitope string

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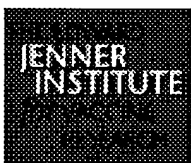
2 Enter the length of amino acid string  
Leave blank to retrieve all.

MIN  MAX

3 Select the restriction allele or serotype

New Search: [MHC Ligand](#) [MHC Kinetics](#) [T Cell Epitope](#) [TCR-MHC](#) [TAP Ligand](#) [Protein L](#)  
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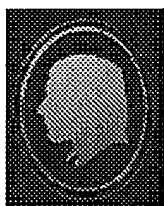
## THE ANTIJEN DATABASE SEARCH RESULTS

You searched for epitope string "ANNPLYKEATSTFTNITYRGT" of undefined length.

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## FULL SEARCH: T CELL EPITOPE

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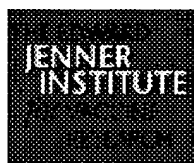
2 Enter the **length of amino acid string**  
Leave blank to retrieve all.

MIN  MAX

3 Select the restriction **allele or serotype**

**New Search:** [MHC Ligand](#) [MHC Kinetics](#) [T Cell Epitope](#) [TCR-MHC](#) [TAP Ligand](#) [Protein I](#)  
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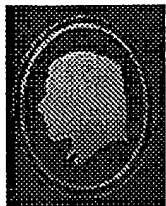
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SEQ ID NO: 17 OF  
6,340,460 / 6,022,544

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## FULL SEARCH: T CELL EPITOPE

1 Enter an epitope string

EPAKSAPAPKKGI

- Replace variable amino acid positions with an asterisk (\*).
- Encase alternative amino acids in square brackets e.g. [AEI]

2 Enter the length of amino acid string

Leave blank to retrieve all.

MIN  MAX

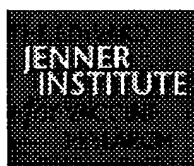
3 Select the restriction allele or serotype

H-2

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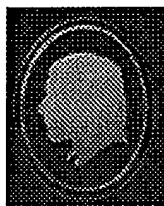
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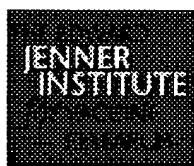
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